

SEQUENCE LISTING

<110> ARES TRADING S.A.

<120> IL-8-LIKE PROTEINS

<130> P032741WO

<140> PCT/GB03/05621

<141> 2003-12-19

<150> GB 0229854.5

<151> 2002-12-20

<160> 24

<170> SeqWin99, version 1.02

<210> 1

<211> 126

<212> DNA

<213> Homo sapiens

<400> 1

atgtcagcac aacatggtct tgtttccaaa tttgggctgg ggcttctgct ccttggggac	60
aaatactttc aaagacatga acaatcaaaa cctcatcaag aagaaataga caacctgcat	120
agccct	126

<210> 2

<211> 42

<212> PRT

<213> Homo sapiens

<400> 2

Met Ser Ala Gln His Gly Leu Val Ser Lys Phe Gly Leu Gly Leu Leu
1 5 10 15

Leu Leu Gly Asp Lys Tyr Phe Gln Arg His Glu Gln Ser Lys Pro His
20 25 30

Gln Glu Glu Ile Asp Asn Leu His Ser Pro
35 40

<210> 3

<211> 330

<212> DNA

<213> Homo sapiens

<400> 3

gacctgccca cgccgggaca cccggtgaca ctccactccc tctgcttttg cagcccccg	60
gggaccctcc tcgagggccc catgtcttct gggttccatc gctttgaggt agaaaatctg	120
aggcctcaaa ctgcccccaa agcaggcaaa ggtcagatgt gtggagagag gatggcgagg	180
atggcaagga cggccaagga gggtcgcccc aggtgcctgg acccagggtt gtcccgacc	240
cgcaccctg gccacatgt ctctcttccc catagcccca ccccagcatc ctggcaccag	300
tgggtcctctg gtggcactgg ctggatgctg	330

<210> 4
 <211> 110
 <212> PRT
 <213> Homo sapiens

<400> 4
 Asp Leu Pro Thr Pro Gly His Pro Val Thr Leu His Ser Leu Cys Phe
 1 5 10 15
 Cys Ser Pro Arg Gly Thr Leu Leu Glu Gly Pro Met Ser Ser Gly Phe
 20 25 30
 His Arg Phe Glu Val Glu Asn Leu Arg Pro Gln Thr Ala Pro Lys Ala
 35 40 45
 Gly Lys Gly Gln Met Cys Gly Glu Arg Met Ala Arg Met Ala Arg Thr
 50 55 60
 Ala Lys Glu Gly Arg Pro Arg Cys Leu Asp Pro Gly Leu Ser Arg Thr
 65 70 75 80
 Pro His Pro Gly Pro His Val Phe Leu Pro His Ser Pro Thr Pro Ala
 85 90 95
 Ser Trp His Gln Trp Ala Pro Gly Gly Thr Gly Trp Met Leu
 100 105 110

<210> 5
 <211> 459
 <212> DNA
 <213> Homo sapiens

<400> 5
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 aaatactttc aaagacatga acaatcaaaa cctcatcaag aagaaataga caacctgcat 120
 agccctgacc tgcccacgcc gggacacccg gtgacactcc actccctctg cttttgcagc 180
 ccccggggga ccctcctcga gggcccatg tcttctgggt tccatcgctt tgaggtagaa 240
 aatctgaggc ctcaaactgc ccccaaagca ggcaaaggtc agatgtgtgg agagaggatg 300
 gcgaggatgg caaggacggc caaggagggt cgcccaggt gcctggaccc aggtttgtcc 360
 cgcaccccg cccctggccc acatgtcttc ctccccata gccccacccc agcatcctgg 420
 caccagtggg ctctggtgg cactggctgg atgctgtag 459

<210> 6
 <211> 152
 <212> PRT
 <213> Homo sapiens

<400> 6
 Met Ser Ala Gln His Gly Leu Val Ser Lys Phe Gly Leu Gly Leu Leu
 1 5 10 15
 Leu Leu Gly Asp Lys Tyr Phe Gln Arg His Glu Gln Ser Lys Pro His
 20 25 30

Gln Glu Glu Ile Asp Asn Leu His Ser Pro Asp Leu Pro Thr Pro Gly
 35 40 45

His Pro Val Thr Leu His Ser Leu Cys Phe Cys Ser Pro Arg Gly Thr
 50 55 60

Leu Leu Glu Gly Pro Met Ser Ser Gly Phe His Arg Phe Glu Val Glu
 65 70 75 80

Asn Leu Arg Pro Gln Thr Ala Pro Lys Ala Gly Lys Gly Gln Met Cys
 85 90 95

Gly Glu Arg Met Ala Arg Met Ala Arg Thr Ala Lys Glu Gly Arg Pro
 100 105 110

Arg Cys Leu Asp Pro Gly Leu Ser Arg Thr Pro His Pro Gly Pro His
 115 120 125

Val Phe Leu Pro His Ser Pro Thr Pro Ala Ser Trp His Gln Trp Ala
 130 135 140

Pro Gly Gly Thr Gly Trp Met Leu
 145 150

<210> 7
 <211> 360
 <212> DNA
 <213> Homo sapiens

<400> 7
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 agaataaaga agttagtaga tggccttgag ttttcccaaa caatggcatt ttctgctacc 120
 aaaataaata tgttattcag tcagaaccac tggactataa gaagtatatt ccattctggt 180
 ttttactggg ggaaaggatg ttgccacaag atgtcagtc atttattcat tcatatatcc 240
 aatagatatt ttatgaccac ttccatgtgc caggagatgg ctaagatcct tggaagacag 300
 ataaaatgct acctaccaac tcaaagtcca gttagggagt cagggggtaa aacaatattc 360

<210> 8
 <211> 120
 <212> PRT
 <213> Homo sapiens

<400> 8
 Asn Thr Glu Asn Asp Phe Tyr Glu Ile Cys Gly Asn Gln Ser His His
 1 5 10 15

His Asp Asn Ala Arg Ile Lys Lys Leu Val Asp Gly Leu Glu Phe Ser
 20 25 30

Gln Thr Met Ala Phe Ser Ala Thr Lys Ile Asn Met Leu Phe Ser Gln
 35 40 45

Asn His Trp Thr Ile Arg Ser Ile Phe His Ser Gly Phe Tyr Trp Gly
 50 55 60

Lys Gly Cys Cys His Lys Met Ser Val His Leu Phe Ile His Ile Ser
65 70 75 80

Asn Arg Tyr Phe Met Thr Thr Ser Met Cys Gln Glu Met Ala Lys Ile
85 90 95

Leu Gly Arg Gln Ile Lys Cys Tyr Leu Pro Thr Gln Ser Pro Val Arg
100 105 110

Glu Ser Gly Gly Lys Thr Ile Phe
115 120

<210> 9
<211> 360
<212> DNA
<213> Homo sapiens

<400> 9
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agaataaaga agttagtaga tggccttgag ttttcccaaa caatggcatt ttctgctacc 120
aaaataaata tgttattcag tcagaaccac tggactataa gaagtatatt ccattctggt 180
ttttactggg ggaaaggatg ttgccacaag atgtcagtc atttattcat tcatatatcc 240
aatagatatt ttatgaccac ttccatgtgc caggagatgg ctaagatcct tggaagacag 300
ataaaatgct acctaccaac tcaaagtcca gttagggagt cagggggtaa aacaatatcc 360

<210> 10
<211> 120
<212> PRT
<213> Homo sapiens

<400> 10
Asn Thr Glu Asn Asp Phe Tyr Glu Ile Cys Gly Asn Gln Ser His His
1 5 10 15

His Asp Asn Ala Arg Ile Lys Lys Leu Val Asp Gly Leu Glu Phe Ser
20 25 30

Gln Thr Met Ala Phe Ser Ala Thr Lys Ile Asn Met Leu Phe Ser Gln
35 40 45

Asn His Trp Thr Ile Arg Ser Ile Phe His Ser Gly Phe Tyr Trp Gly
50 55 60

Lys Gly Cys Cys His Lys Met Ser Val His Leu Phe Ile His Ile Ser
65 70 75 80

Asn Arg Tyr Phe Met Thr Thr Ser Met Cys Gln Glu Met Ala Lys Ile
85 90 95

Leu Gly Arg Gln Ile Lys Cys Tyr Leu Pro Thr Gln Ser Pro Val Arg
100 105 110

Glu Ser Gly Gly Lys Thr Ile Phe
115 120

<210> 11
 <211> 265
 <212> DNA
 <213> Homo sapiens

<400> 11
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 agtatattcc attctgggtt ttactggggg aaaggatgtt gccacaagat gtcagtccat 120
 ttattcattc atatatccaa tagatatattt atgaccactt ccatgtgcca ggagatggct 180
 aagatccttg gaagacagat aaaatgctac ctaccaactc aaagtccagt tagggagtca 240
 gggggtaaaa caatattcta gcaca 265

<210> 12
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 12
 Met Ala Phe Ser Ala Thr Lys Ile Asn Met Leu Phe Ser Gln Asn His
 1 5 10 15
 Trp Thr Ile Arg Ser Ile Phe His Ser Gly Phe Tyr Trp Gly Lys Gly
 20 25 30
 Cys Cys His Lys Met Ser Val His Leu Phe Ile His Ile Ser Asn Arg
 35 40 45
 Tyr Phe Met Thr Thr Ser Met Cys Gln Glu Met Ala Lys Ile Leu Gly
 50 55 60
 Arg Gln Ile Lys Cys Tyr Leu Pro Thr Gln Ser Pro Val Arg Glu Ser
 65 70 75 80
 Gly Gly Lys Thr Ile Phe
 85

<210> 13
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer INSP094-CP1

<400> 13
 atggcatttt ctgctaccaa 20

<210> 14
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer INSP094-CP2

<400> 14
tgtgctagaa tattgtttta cc 22

<210> 15
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer INSP094-EX1

<400> 15
aagcaggctt cgccaccatg gcatttttctg ctaccaa 37

<210> 16
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer INSP094-EX2

<400> 16
gtgatgggga ttggtggaata ttgttttacc ccctg 35

<210> 17
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer GCP Forward

<400> 17
ggggacaagt ttgtacaaaa aagcaggctt cgccacc 37

<210> 18
<211> 51
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer GCP Reverse

<400> 18
ggggaccact ttgtacaaga aagctggggt tcaatgggtga tggatgatgg g 51

<210> 19
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer pEAK12F

<400> 19
gccagcttgg cacttgatgt 20

<210> 20
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer pEAK12R

<400> 20
gatggaggtg gacgtgtcag 20

<210> 21
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer 21M13

<400> 21
tgtaaaacga cggccagt 18

<210> 22
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer M13REV

<400> 22
caggaaacag ctatgacc 18

<210> 23
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer T7

<400> 23
taatacgact cactatagg 19

<210> 24
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer T3

<400> 24
attaaccctc actaaagg

18